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Sequence Listing was accepted.
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Reviewer: Anne Corrigan
Timestamp: [year=2008; month=9; day=29; hr=9; min=44; sec=3; ms=408;]
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Validated By CRFValidator v 1.0.3

Application No: 10561720 Version No: 2.0

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Finished: 2008-08-29 16:36:15.776
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Total Errors: 0
No. of SeqIDs Defined: 19
Actual SeqID Count: 19

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<110> Board of Trustees Operating Michigan State University
Allison, Richard F.

<120> Expression of a Recombinant Transgene

<130> 6550-000072/US/NPB

<140> 10561720
<141> 2005-12-22

<150> PCT/US04/21451
<151> 2004-07-02

<150> US 60/485,073
<151> 2003-07-03

<160> 19

<170> PatentIn version 3.5

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<212> DNA
<213> Cowpea chlorotic mottle virus

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<210> 2
<211> 16
<212> DNA
<213> Cowpea chlorotic mottle virus

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<212> DNA
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aagatatatt tctcaagatc agaagtacta ttccagtatg gacgattcaa ggcttgcttc 240

acaaaccaag gcaagtaata gagattggag tctctaaaaa ggtagttccc actgaatcaa 300

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tggagcacga cacacttgtc tactccaaaa atatcaaaga tacagtctca gaagaccaaa 480

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<210> 4
<211> 581
<212> DNA
<213> Encephalomyocarditis virus

<400> 4
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gggcccggaa acctggccct gtcttcttga cgagcattcc taggggtctt tcccctctcg 180

ccaaaggaat gcaaggtctg ttgaatgtcg tgaaggaagc agttcctctg gaagcttctt 240

gaagacaaac aacgtctgta gcgacccttt gcaggcagcg gaaccccca cctggcgaca 300

ggtgcctctg cggccaaaag ccacgtgtat aagatacacc tgcaaaggcg gcacaacccc 360

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tcaacaaggg gctgaaggat gcccagaagg taccccattg tatgggatct gatctggggc 480

ctcggtgcac atgctttaca tgtgtttagt cgaggttaaa aaaacgtcta ggccccccga 540

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<210> 5
<211> 581
<212> RNA
<213> Encephalomyocarditis virus

<400> 5
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gggcccggaa accuggcccu gucuucuuga cgagcauucc uaggggucuu uccccucucg 180

ccaaaggaau gcaaggucug uugaaugucg ugaaggaagc aguuccucug gaagcuucuu 240

gaagacaaac aacgucugua gcgacccuuu gcaggcagcg gaacccccca ccuggcgaca 300

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ucaacaaggg gcugaaggau gcccagaagg uaccccauug uaugggaucu gaucuggggc 480

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<212> DNA
<213> Encephalomyocarditis virus

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acaatggggt accttctggg catccttcag ccccttgttg aatacgcttg aggagagcca 180

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<210> 7
<211> 581
<212> RNA
<213> Encephalomyocarditis virus

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acaauggggu accuucuggg cauccuucag ccccuuguug aauacgcuug aggagagcca 180

uuugacucuu uccacaacua uccaacucac aacguggcac ugggguugug ccgccuuugc 240

agguguaucu uauacacgug gcuuuuggcc gcagaggcac cugucgccag gugggggguu 300

ccgcugccug caaagggucg cuacagacgu uguuugucuu caagaagcuu ccagaggaac 360

ugcuuccuuc acgacauuca acagaccuug cauuccuuug gcgagagggg aaagaccccu 420

aggaaugcuc gucaagaaga cagggccagg uuuccgggcc cucacauugc caaaagacgg 480

caauauggug gaaaaucaca uauagacaaa cgcacaccgg ccuuauucca agcggcuucg 540

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<210> 8
<211> 242
<212> DNA
<213> Cowpea chlorotic mottle virus

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ggttttactc cttgaaccct tcggaagaac tctttggagt tcgtaccagt acctcacata 180

gtgaggtaat aagactggtg ggcagcgcct agtcgaaaga ctaggtgatc tctaaggaga 240

cc 242

<210> 9
<211> 242
<212> RNA
<213> Cowpea chlorotic mottle virus

<400> 9
agugcccgcu gaagagcguu acacuagugu ggccuacuug aaggcuaguu auaaccguuu 60

cuuuaaacgg uaaucguugu ugaaacgucu uccuuuuaca agaggauuga gcugcccuug 120

gguuuuacuc cuugaacccu acggaagaac ucuuuggagu ucguaccagu accucacaua 180

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cc 242

<210> 10
<211> 242
<212> DNA
<213> Cowpea chlorotic mottle virus

<400> 10
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actatgtgag gtactggtac gaactccaaa gagttcttcc gaagggttca aggagtaaaa 120

cccaagggca gctcaatcct cttgtaaaag gaagacgttt caacaacgat taccgtttaa 180

agaaacggtt ataactagcc ttcaagtagg ccacactagt gtaacgctct tcagcgggca 240

ct 242

<210> 11
<211> 242
<212> RNA
<213> Cowpea chlorotic mottle virus

<400> 11
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acuaugugag guacugguac gaacuccaaa gaguucuucc gaaggguuca aggaguaaaa 120

cccaagggca gcucaauccu cuuguaaaag gaagacguuu caacaacgau uaccguuuaa 180

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cu 242

<210> 12
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<212> DNA
<213> Artificial

<220>
<223> Artificial sequence used to show antisense relationship of a gene and IRES to a promoter and viral 3' UTR

<220>
<221> misc_feature
<222> (1)..(3)
<223> n is a, c, g, or t

<400> 12
nnncatggaa tt 12

<210> 13
<211> 12
<212> DNA
<213> Artificial

<220>
<223> Complement of artificial sequence used to show antisense relationship of a gene and IRES to a promoter and viral 3' UTR

<220>
<221> misc_feature
<222> (10)..(12)
<223> n is a, c, g, or t

<400> 13
aattccatgn nn 12

<210> 14
<211> 12
<212> RNA
<213> Artificial

<220>
<223> Transcript of RNA polymerase

<220>
<221> misc_feature
<222> (1)..(3)
<223> n is a, c, g, or u

<400> 14
nnncauggaa uu 12

<210> 15
<211> 12
<212> RNA
<213> Artificial

<220>
<223> Complement of transcript of RNA polymerase

<220>
<221> misc_feature
<222> (10)..(12)
<223> n is a, c, g, or u

<400> 15
aauuccaugn nn 12

<210> 16
<211> 12
<212> DNA
<213> Artificial Sequence

<220>
<223> DNA Construct containing promoter complementary coding sequence, exemplary IRES complementary sequence and a viral 3' UTR in 5' - 3' orintation

<220>
<221> misc_feature
<223> DNA construct wherein YYY indicates complementary first translatable codon after initiation codon and an asterisk indicates a stop codon.

<400> 16
yyycatggaa tt 12

<210> 17
<211> 12
<212> DNA
<213> Artificial Sequence

<220>
<223> DNA Construct containing promoter, coding sequence, exemplary IRES sequence and a viral 3' UTR in 3' - 5' orintation

<220>
<221> misc_feature
<223> DNA construct wherein XXX indicates first translatable codon after initiation codon and an asterisk indicates a stop codon.

<400> 17
yyygtacctt aa 12

<210> 18
<211> 12
<212> RNA
<213> Artificial Sequence

<220>
<223> RNA Construct containing complementary coding sequence, exemplary IRES complementary sequence and a viral 3' UTR in 5' - 3' orintation

<220>
<221> misc_feature
<223> Recombinant RNA sequence where YYY is the complement of the first codon after the initiation codon and where an asterisk indicates a stop codon.

<400> 18
yyycauggaa uu 12

<210> 19
<211> 12
<212> RNA
<213> Artificial Sequence

<220>
<223> RNA Construct containing viral 3' UTR, exemplary IRES sequence and a coding sequence in 5' - 3' orientation

<220>
<221> misc_feature

<223> Complementary sequence (sense strand) of RNA recombinant sequence
where XXX is the first translatable codon after initiation codon
and where an asterisk indicates a stop codon.

<400> 19
aauuccaugy yy 12